

# Why the *in silico* Drug Design and Discovery

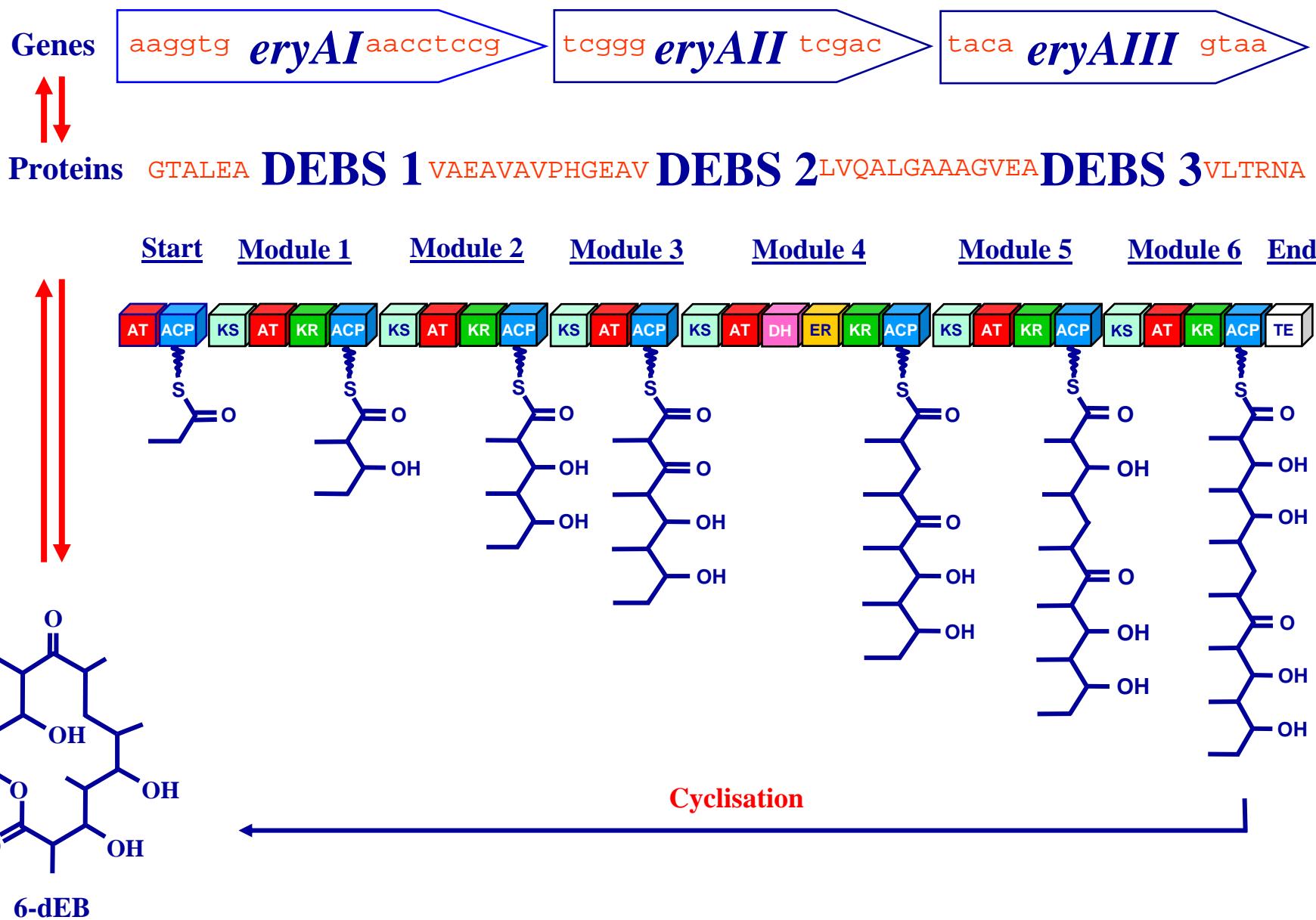


*and*

**Novalis**

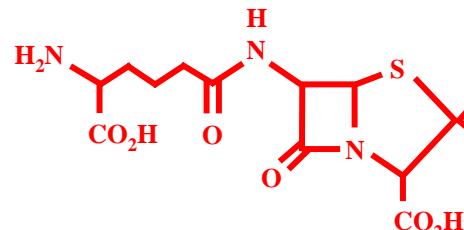
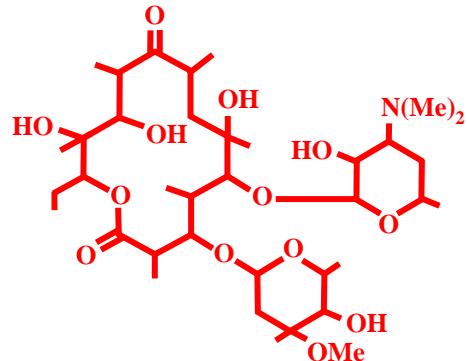
**Daslav Hranueli**

# *Modular type I PKS, e.g. erythromycin:*



# *Biological activity of polyketides and peptides:*

*Erythromycin A*

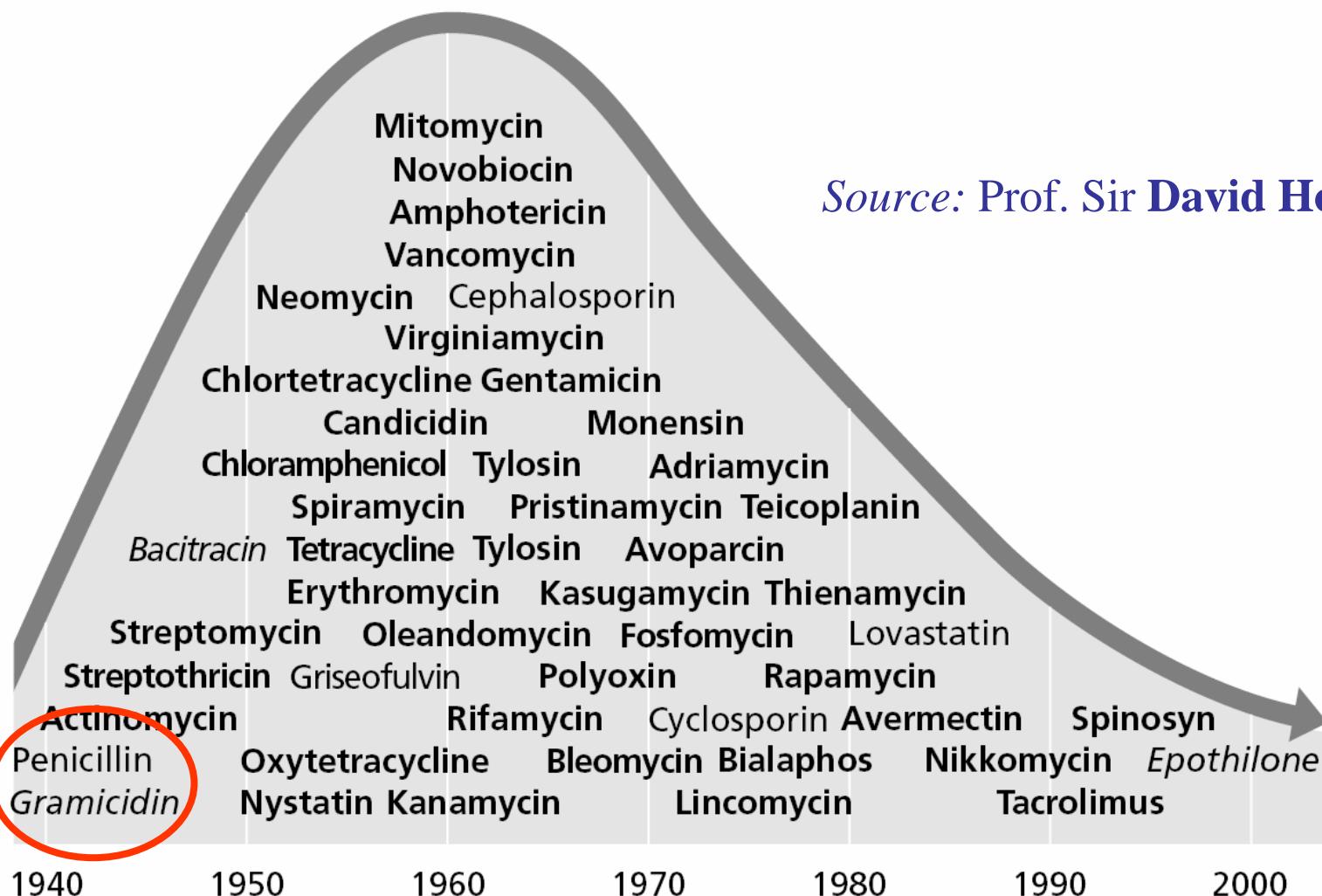


*Isopenicillin N*

**50% of most important medications:**

- not only
  - antibacterials,
  - antifungals,
  - antivirals, and
  - antitumor compounds
- but also products with
  - immunosuppressants,
  - antihypertensives,
  - antidiabetics,
  - antimalarials, and
  - antiosteoporotics

# *Biological activity of polyketides and peptides:*



Source: Prof. Sir David Hopwood

# *Are there more Actinomycetes to be discovered?*

- New Species - PubMed search:

"Actinomycete sp. nov." (30/05/08)	-	2006	-	40
	-	2007	-	38
	-	2008	-	18

- New Genera - PubMed search:

"Actinomycete gen. nov. sp. nov." (30/05/08)	-	2006	-	7
	-	2007	-	4
	-	2008	-	2

(Richard Baltz, Cubist Pharmaceuticals, Inc., USA

Natural products discovery & Production II, Whistler, British Columbia, Canada, 2008)

*Biological activity of polyketides and peptides:*

*Alternative strategies*

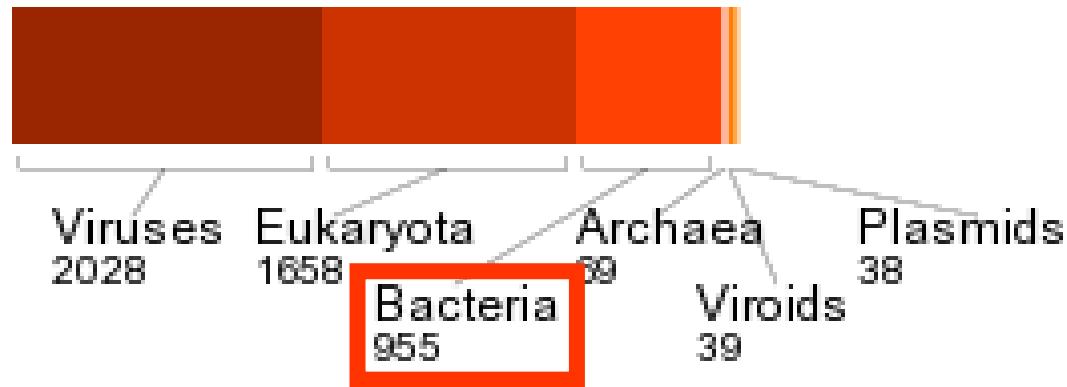
*to explore chemical potential*

*of natural products?*

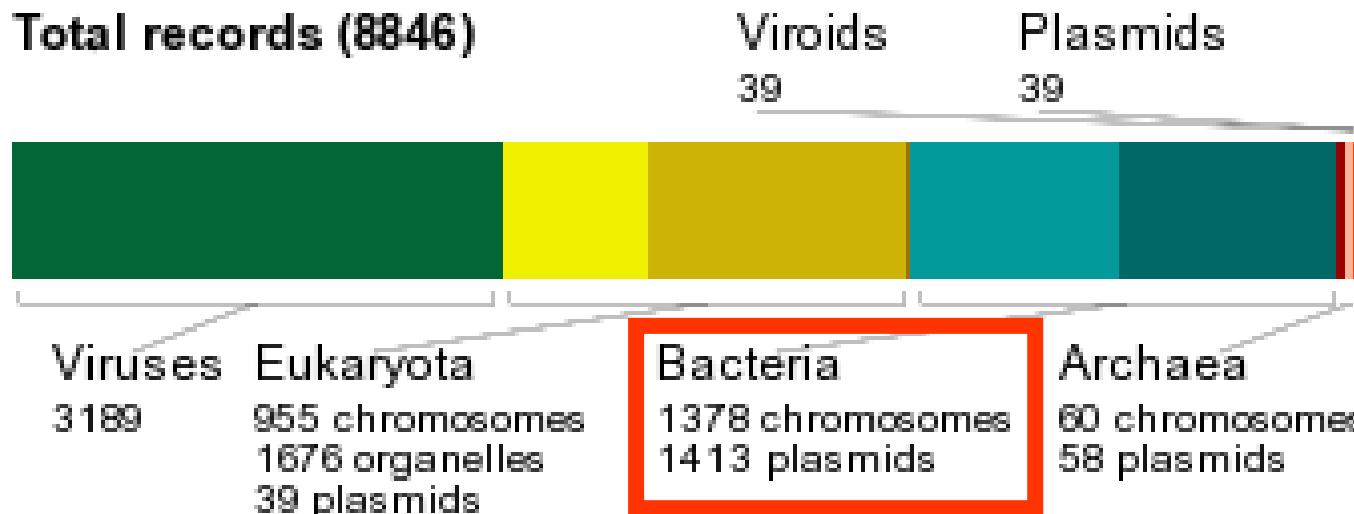
# *Current Status of Genome Sequencing Projects*

## NCBI (16/10/2008)

Total species (4787)



Total records (8846)



# *PKS and NRPS gene-clusters in bacteria:*

Species	Size (MB)	PKS/NRPS (KB)
<i>Zymomonas mobilis</i>	2.0	0
<i>Gluconobacter oxydans</i>	2.9	0
<i>Clostridium perfringens</i>	3.1	0
<i>Brucella abortis</i>	3.3	3
<i>Corynebacterium glutamicum</i>	3.3	0
<i>Bacillus licheniformis</i>	4.2	35
<i>Escherichia coli K12</i>	4.7	5.5
<i>Xanthomonas campestris</i>	5.1	4
<i>Salinospora tropica</i>	5.2	516
<i>Photorhabdis luminescens</i>	5.7	208
<i>Pseudomonas aeruginosa</i>	6.3	74
<i>Nocardia farcinica</i>	6.3	204
<i>Pseudomonas fluorescens</i>	7.1	183
<i>Saccharopolyspora erythraea</i>	8.2	410
<i>Streptomyces coelicolor</i>	8.6	135
<i>Streptomyces avermitilis</i>	9.2	356

(Donadio *et al.*, *Nat. Prod. Rep.*, **24**, 1073, 2007)

# PKS and NRPS gene-clusters cont':

Secondary metabolite - Mozilla

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EBI Databases Nucleotide for Go Clear

13	polyketide	pks11	113361	118594	2	5234
14	polyketide	pte	487415	567017	13	79603
15	polyketide	ave	1132451	1012000		80916
16	polyketide	pks2	1895794	1913284	4	17491
17	polyketide	pks3	2773878	2784841	7	10964
18	polyketide	pks5	2877941	2894413	4	16473
19	polyketide	pks9	2897395	2913535	15	16141
20	polyketide	olm	353561	3624500	11	100068
21	polyketide	pks8	4527901	4540450	14	12550
22	polyketide	rpp	8490207	8492484	2	2278
23	polyketide	pks4	8553602	8561604	3	8003
24	polyketide	pks1	8776963	8789766	4	12804
			H <sub>2</sub> N			
			N			
			N			
			H			
			Subtotal		OH	98
						362525
25	peptide	nrps6	754376	763274	7	8899
26	peptide	nrps7	991484	1042269	32	50786
27	peptide	nrps8	1549424	1554224	3	4801
28	peptide	nrps3	3930088	3938730	6	8643
29	peptide	nrps1	3974805	3996189	7	21385
30	peptide	nrps2	4494250	4526990	16	32741
31	peptide	nrps5	7930737	7937201	4	6465
32	peptide	nrps4	8522760	8530697	5	7938
					Subtotal	80
						141658

11 PKS gene-clusters

Avermectin

S. coelicolor 2 PKS, 4 NRPS and 2 PKS/NRPS

Sac. erythrea 11 PKS, 6 NRPS and 1 PKS/NRPS

S. griseus 6 PKS, 7 NRPS and 2 PKS/NRPS

S. scabies 3 PKS, 4 NRPS and 1 PKS/NRPS

8 NRPS gene-clusters

Total 33 PKS, 29 NRPS and 6 PKS/NRPS

Start

Genome P... Kongres - ... Kalendar - ... Removabl... Secondar...

14:10

68 gene-clusters in only 5 genomes

# Current Status of Genome Sequencing Projects

## NCBI (16/10/2008) cont'

Microbial Genome Sequencing Projects in Progress - Microsoft Internet Explorer

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Address http://www.ncbi.nlm.nih.gov/genomes/lproks.cgi Go Links

<a href="#">352 Streptococcus suis P1/7</a>	B	Firmicutes	-	-	<b>*1.83</b>	<b>40</b>	-	-	Sanger Insti	
<a href="#">353 Streptococcus uberis 0140T</a>	B	Firmicutes	-	-	<b>*1.82</b>	<b>40</b>	-	-	Sanger Insti	
<a href="#">93 Streptomyces ambofaciens ATCC 23877</a>	B	Actinobacteria	-	-	<b>*8</b>	<b>72</b>	-	-	Genoscope	
<a href="#">19249 Streptomyces clavuligerus ATCC 27064</a>	B	Actinobacteria	-	-	-	-	-	-	Korea Resear	
<a href="#">28551 Streptomyces clavuligerus ATCC 27064</a>	B	Actinobacteria	<b>597</b>	<b>5981</b>	-	<b>71.1</b>	<a href="#">NZ_ABH00000000</a>	<a href="#">BLAST</a>	<b>02/28/08</b>	Broad Instit
<a href="#">13658 Streptomyces peucetius ATCC 27952</a>	B	Actinobacteria	-	-	-	-	-	-	Institute of	
<a href="#">28553 Streptomyces pristinaespiralis ATCC 25486</a>	B	Actinobacteria	<b>532</b>	<b>5591</b>	-	<b>70.4</b>	<a href="#">NZ_ABI00000000</a>	<a href="#">BLAST</a>	<b>02/28/08</b>	Broad Instit
<a href="#">12985 Streptomyces scabiei</a>	B	Actinobacteria	-	-	-	-	-	-	Sanger Insti	
<a href="#">28547 Streptomyces sp. Mg1</a>	B	Actinobacteria	<b>466</b>	<b>6637</b>	-	<b>71.2</b>	<a href="#">NZ_ABF00000000</a>	<a href="#">BLAST</a>	<b>02/28/08</b>	Broad Instit
<a href="#">28549 Streptomyces sp. SPB74</a>	B	Actinobacteria	<b>534</b>	<b>4638</b>	-	<b>71.9</b>	<a href="#">NZ_ABG00000000</a>	<a href="#">BLAST</a>	<b>02/28/08</b>	Broad Instit
<a href="#">28555 Streptomyces sviceus ATCC 29083</a>	B	Actinobacteria	<b>522</b>	<b>7990</b>	-	<b>70.1</b>	<a href="#">NZ_ABJ00000000</a>	<a href="#">BLAST</a>	<b>02/28/08</b>	Broad Instit
<a href="#">21083 Streptosporangium roseum DSM 43021</a>	B	Actinobacteria	-	-	-	<b>69-</b>	<b>71</b>	-	-	DOE Joint
<a href="#">18181 Subdoligranulum variabile DSM 15176</a>	B	Firmicutes	-	-	-	<b>52.2</b>	-	-	-	Genome Ins
										Washington
										University
										(WashU)

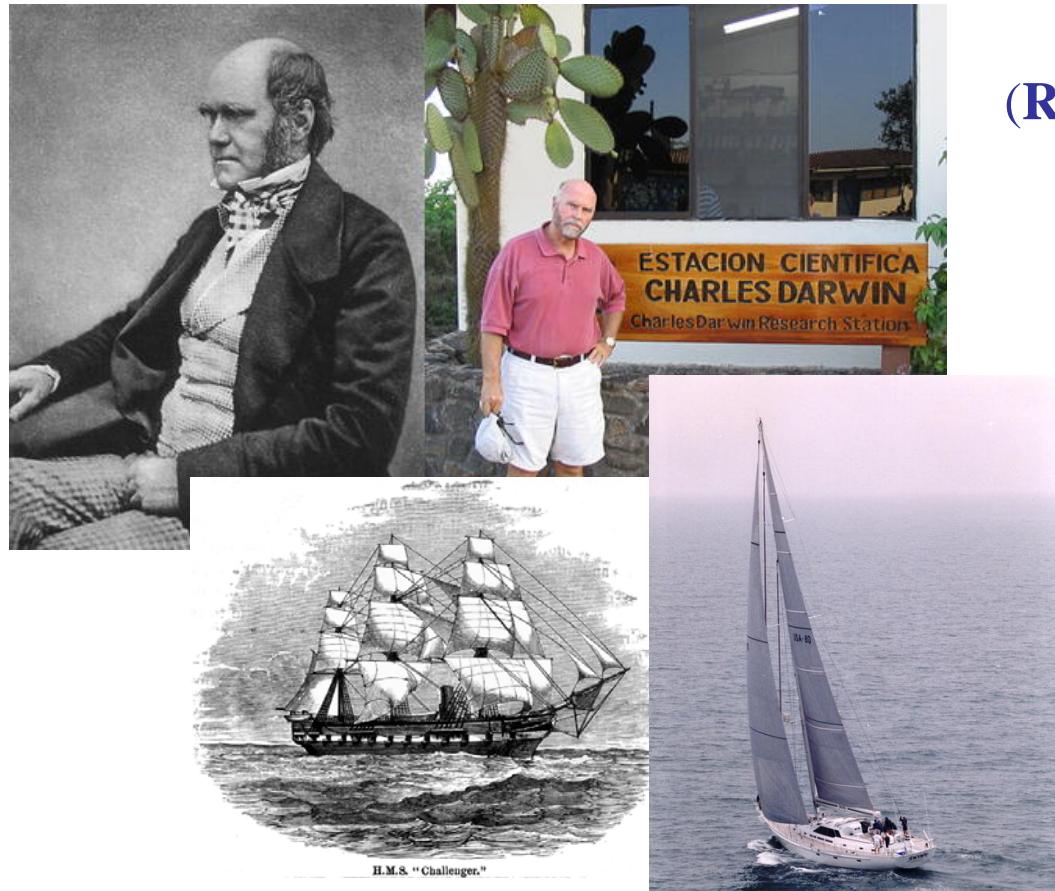
10 Actinobacteria

*PKS and NRPS gene-clusters cont':*

*Significant chemical diversity*

*potential in sequenced genomes*

# *Sequencing of largest metagenomic dataset:*



(Rusch *et al.*, PLoS Biol. 5, e77, 2007)

## CAMERA database

- **7.7 million sequencing reads**
- **6.3 billion bp of DNA**

*Sequencing of largest metagenomic dataset:*

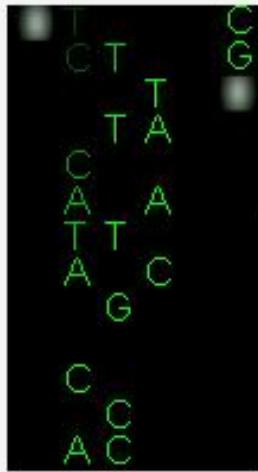
*Computational tools needed for rapid  
mining of massive DNA datasets to  
hunt for gene clusters encoding  
entirely novel compounds*

# *Program packages:*



*and* **Novalis**

Version 1.00 beta1



*Cluster Scanner*

**Java and JavaScript; Linux server (<http://bioserv.pbf.hr/>)**

**Java client on user's computer**